

What is Teleost Taxonomy?

The Teleost taxonomy ontology is being used to facilitate annotation of phenotypes, particularly for taxa that are not covered by NCBI because no submissions of molecular data have been made. Taxonomy ontologies can also be valuable in annotating legacy data, where authors make phenotype or ecological assertions (e.g., host-parasite associations) that refer to groups that are reorganized or no longer recognized. The taxonomy ontology serves as the source of taxa for our project's use for identifying evolutionary changes that match the phenotype of a zebrafish mutant.

Discuss Teleost Taxonomy Terminology

Please go to the [Teleost Taxonomy Discussion](#) page to leave comments regarding the Teleost Taxonomy Terminology.

Examples of comments may include:

- How you are using this terminology
- Why you decided to use (or not use) this terminology
- Strengths or limitations of the terminology
- Comparisons to other similar terminologies

Links

[Teleost Taxonomy FAQ](#)

[Teleost Taxonomy Home Page](#)